

#



DECLARATION

In re application of Pierre Legrain and al., serial number N° 10/023,530 filed on December 18th, 2001, for PROTEINS THAT INTERACT WITH β TrCP :

I prepared the electronic version and the paper copy of the Sequence Listing in connection with the captioned US patent application. The Sequence Listing information recorded in the computer readable form is identical to the written paper Sequence Listing.

A handwritten signature in black ink, appearing to read "Aude Guenard", written over a horizontal line.

Aude GUENARD

#4



B4717A-US.ST25.txt
SEQUENCE LISTING

<110> LEGRAIN, Pierre
BENAROUS, Richard
BLOT, Guillaume
LASSOT, Irina

<120> PROTEINS THAT INTERACT WITH BETA TrCP

<130> B4717A

<140> 10/023,530

<141> 2001-12-18

<150> 60/256,276

<151> 2000-12-18

<160> 30

<170> PatentIn version 3.1

<210> 1

<211> 657

<212> DNA

<213> Homo sapiens

<220>

<221> gene

<222> (1)..(657)

<223> Beta TrCP

<400> 1
atggaccg ccgaggcggt gctgcaagag aaggcactca agtttatgaa ttcctcagag 60
agagaagact gtaataatgg cgaacccccct aggaagataa taccagagaa gaattcactt 120
agacagacat acaacagctg tgccagactc tgcttaaacc aagaaacagt atgttttagca 180
agcactgcta tgaagactga gaattgtgtg gccaaaacaa aacttgccaa tggcacttcc 240
agtatgattg tgcccaagca acggaaactc tcagcaagct atgaaaagga aaaggaactg 300
tgtgtcaa atctttgagca gtggtcagag tcagatcaag tggaatttgt ggaacatctt 360
atatcccaaa tgtgtcatta ccaacatggg cacataaact cgtatcttaa acctatgttg 420

B4717A-US.ST25.txt

cagagagatt tcataactgc tctgccagct cggggattgg atcatatcgc tgagaacatt 480
 ctgtcatacc tggatgcaa atcactatgt gctgctgaac ttgtgtgcaa ggaatggtag 540
 cgagtgcact ctgatggcat gctgtggaag aagcttatcg agagaatggg caggacagat 600
 tctctgtgga gaggcctggc agaacgaaga ggatggggac agtatttatt caaaaac 657

<210> 2

<211> 219

<212> PRT

<213> Homo sapiens

<220>

<221> BetaTrCP

<222> (1)..(219)

<223> F-box protein

<400> 2

Met Asp Pro Ala Glu Ala Val Leu Gln Glu Lys Ala Leu Lys Phe Met
 1 5 10 15

Asn Ser Ser Glu Arg Glu Asp Cys Asn Asn Gly Glu Pro Pro Arg Lys
 20 25 30

Ile Ile Pro Glu Lys Asn Ser Leu Arg Gln Thr Tyr Asn Ser Cys Ala
 35 40 45

Arg Leu Cys Leu Asn Gln Glu Thr Val Cys Leu Ala Ser Thr Ala Met
 50 55 60

Lys Thr Glu Asn Cys Val Ala Lys Thr Lys Leu Ala Asn Gly Thr Ser
 65 70 75 80

Ser Met Ile Val Pro Lys Gln Arg Lys Leu Ser Ala Ser Tyr Glu Lys
 85 90 95

Glu Lys Glu Leu Cys Val Lys Tyr Phe Glu Gln Trp Ser Glu Ser Asp
 100 105 110

Gln val Glu Phe Val Glu His Leu Ile Ser Gln Met Cys His Tyr Gln
 115 120 125

His Gly His Ile Asn Ser Tyr Leu Lys Pro Met Leu Gln Arg Asp Phe
 130 135 140

B4717A-US.ST25.txt

Ile Thr Ala Leu Pro Ala Arg Gly Leu Asp His Ile Ala Glu Asn Ile
145 150 155 160

Leu Ser Tyr Leu Asp Ala Lys Ser Leu Cys Ala Ala Glu Leu Val Cys
165 170 175

Lys Glu Trp Tyr Arg Val Thr Ser Asp Gly Met Leu Trp Lys Lys Leu
180 185 190

Ile Glu Arg Met Val Arg Thr Asp Ser Leu Trp Arg Gly Leu Ala Glu
195 200 205

Arg Arg Gly Trp Gly Gln Tyr Leu Phe Lys Asn
210 215

<210> 3

<211> 1680

<212> DNA

<213> Homo sapiens

<220>

<221> gene

<222> (1)..(1680)

<223> Ras SF1

<400> 3
ccggggcggt gggtggcggc tacggacgcg caggactggg ggacgggagg gtacggctat 60
gggagaggcg gaggcgcctt ctttcgaaat gacctggagc agcacgacga gcagtggcta 120
ctgcagccaa gaggactcgg actcggagct cgagcagtag ttcaccgagc gaacctcgtc 180
agctcgcagg ccgcgcccgg accaggacga gcctgtggag tgggagacac ctgacctttc 240
tcaagctgag attgagcaga agatcaagga gtacaatgcc cagatcaaca gcaacctctt 300
catgagcttg aacaaggacg gttcttacac aggcttcac aagggttcagc tgaagctggg 360
gcgccctgtc tctgtgccct ccagcaagaa gccaccctcc ttgcaggatg cccggcgggg 420
cccaggacgg ggcacaagtg tcaggcgccg cacttccttt tacctgcca aggatgctgt 480
caagcacctg catgtgctgt cacgcacaag ggcacgtgaa gtcattgagg ccctgctgag 540
aaagttcttg gtggtggatg acccccgcaa gtttgactc tttgagcgcg ctgagcgtca 600
cggccaagtg tacttgcgga agctgttga tgatgagcag cccctgcggc tgcggctcct 660
ggcagggccc agtgacaagg ccctgagctt tgcctgaag gaaaatgact ctggggaggt 720
gaactgggac gccttcagca tgcctgaact acataacttc ctacgtatcc tgcagcggga 780

B4717A-US.ST25.txt

```

ggaggaggag cacctccgcc agatcctgca gaagtactcc tattgccgcc agaagatcca      840
agaggccctg cagcctgcc cccttgggtg acctcttgta ccccagggtg gaaggcagac      900
agcaggcagc gccaaagtgcg tgccgtgtga gtgtgacagg gccagtgggg cctgtggaat      960
gagtgtgcat ggaggccctc ctgtgctggg ggaatgagcc cagagaacag cgaagtagct     1020
tgctccctgt gtccacctat ggggtgtagc aggtatggct ctgcaccctt ctgccctcat     1080
tactgggcct tagtgggcca gggctgccct gagaagctgc tccaggcctg cagcaggagt     1140
ggtgcagaca gaagtctcct caatttttgt ctcagaagtg aaaatcttgg agaccctgca     1200
aacagaacag ggtcatgttt gcaggggtga cggccctcat ctatgaggaa aggttttgga     1260
tcttgaatgt ggtctcagga tatccttatc agagctaagg gtgggtgctc agaataaggc     1320
aggcattgag gaagagtctt ggtttctctc tacagtgcc aactctcaca caccctgagg     1380
tcaggggagt ctggctcaca gtacagcatg tgccttaatg cttcatatga ggaggatgtc     1440
cctgggccag ggtctgtgtg aatgtgggca ctggcccagg ttcatacctt atttgctaata     1500
caaagccagg gtctctccct cagggtgttt ttatgaagtg cgtgaatgta tgtaatgtgt     1560
ggtggcctca gctgaatgcc tcctgtgggg aaaggggttg gggtgacagt catcatcagg     1620
cctggggctg agagaattgg ctcaataaag atttcaagat ccaaaaaaaaa aaaaaaaaaa     1680

```

<210> 4

<211> 270

<212> PRT

<213> Homo sapiens

<220>

<221> RasSF1

<222> (1)..(270)

<223> tumor suppressor

<400> 4

Met Gly Glu Ala Glu Ala Pro Ser Phe Glu Met Thr Trp Ser Ser Thr
1 5 10 15

Thr Ser Ser Gly Tyr Cys Ser Gln Glu Asp Ser Asp Ser Glu Leu Glu
20 25 30

Gln Tyr Phe Thr Ala Arg Thr Ser Leu Ala Arg Arg Pro Arg Arg Asp
35 40 45

Gln Asp Glu Pro Val Glu Trp Glu Thr Pro Asp Leu Ser Gln Ala Glu
50 55 60

B4717A-US.ST25.txt

Ile Glu Gln Lys Ile Lys Glu Tyr Asn Ala Gln Ile Asn Ser Asn Leu
65 70 75 80

Phe Met Ser Leu Asn Lys Asp Gly Ser Tyr Thr Gly Phe Ile Lys Val
85 90 95

Gln Leu Lys Leu Val Arg Pro Val Ser Val Pro Ser Ser Lys Lys Pro
100 105 110

Pro Ser Leu Gln Asp Ala Arg Arg Gly Pro Gly Arg Gly Thr Ser Val
115 120 125

Arg Arg Arg Thr Ser Phe Tyr Leu Pro Lys Asp Ala Val Lys His Leu
130 135 140

His Val Leu Ser Arg Thr Arg Ala Arg Glu Val Ile Glu Ala Leu Leu
145 150 155 160

Arg Lys Phe Leu Val Val Asp Asp Pro Arg Lys Phe Ala Leu Phe Glu
165 170 175

Arg Ala Glu Arg His Gly Gln Val Tyr Leu Arg Lys Leu Leu Asp Asp
180 185 190

Glu Gln Pro Leu Arg Leu Arg Leu Leu Ala Gly Pro Ser Asp Lys Ala
195 200 205

Leu Ser Phe Val Leu Lys Glu Asn Asp Ser Gly Glu Val Asn Trp Asp
210 215 220

Ala Phe Ser Met Pro Glu Leu His Asn Phe Leu Arg Ile Leu Gln Arg
225 230 235 240

Glu Glu Glu Glu His Leu Arg Gln Ile Leu Gln Lys Tyr Ser Tyr Cys
245 250 255

Arg Gln Lys Ile Gln Glu Ala Leu His Ala Cys Pro Leu Gly
260 265 270

<210> 5

<211> 10

<212> DNA

<213> Artificial sequence

<220>

<223> linker to clone into pP6 plasmid

<220>

<221> HGX931

<222> (1)..(10)

<223>

<400> 5
gggccacgaa

10

<210> 6

<211> 13

<212> DNA

<213> Artificial sequence

<220>

<223> linker to clone into pP6 plasmid

<220>

<221> HGX932

<222> (1)..(13)

<223>

<400> 6
ttcgtggccc ctg

13

<210> 7

<211> 69

<212> DNA

<213> Artificial sequence

<220>

<223> oligonucleotide used for the construction of pP6 plasmid

<220>

<221> oligonucleotide

<222> (1)..(69)

<223> 5'-3' single-stranded
complementary to sequence ID NO.28

<400> 7

tcgagggggc cccagtggcc ctttaattaa ggatccccac tagtgcggcc gcggcccctg 60

cggccatgg 69

<210> 8

<211> 64

<212> DNA

<213> Artificial Sequence

<220>

<223> oligonucleotide used for the construction of pB6 plasmid

<220>

<221> oligonucleotide

<222> (1)..(64)

<223> 5'-3' single-stranded
Complementary to sequence SEQ ID NO. 29

<400> 8
tcgagggggc cccagtggcc ctttaattaa ggatccccac tagtgcggcc gcggcccgtc 60

cggc 64

<210> 9

<211> 68

<212> DNA

<213> Artificial Sequence

<220>

<223> oligonucleotide used for the construction of pB20 plasmid

<220>

<221> oligonucleotide

<222> (1)..(68)

<223> 5'-3' single-stranded
Complementary to SEQ ID NO. 30

<400> 9
ggtcgagggg cccagtggc ctttaattaa ggatccccac tagtgcggcc gcggcccgtc 60

cggccccg 68

<210> 10
 <211> 20
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> primer used in PCR on yeast colonies
 <220>
 <221> ABS1
 <222> (1)..(20)
 <223> PCR primer

<400> 10
 gcgtttggaa tcactacagg

20

<210> 11
 <211> 19
 <212> DNA
 <213> Artificial Sequence

<220>
 <223> Primer used in PCR on yeast colonies
 <220>
 <221> ABS2
 <222> (1)..(19)
 <223> PCR primer

<400> 11
 cacgatgcac gttgaagtg

19

<210> 12
 <211> 36
 <212> DNA
 <213> homo sapiens

<220>

<221> TrCP(1-260)sense

<222> (1)..(36)

<223> Beta-TrCP primer

<400> 12

gcgagcggat ccatggaccc ggccgaggcg gtgctg

36

<210> 13

<211> 42

<212> DNA

<213> homo sapiens

<220>

<221> TrCP(1-260)antisense

<222> (1)..(42)

<223> Beta-TrCP primer

<400> 13

gcgagcgtcg acctagggcc ctctctgtaa actatgtctt cc

42

<210> 14

<211> 39

<212> DNA

<213> homo sapiens

<220>

<221> TrCP(261-569)sense

<222> (1)..(39)

<223> Beta-TrCP primer

<400> 14

gcgagcggat ccatgattca ctgccgaagt gaaacaagc

39

<210> 15

<211> 42

<212> DNA

<213> homo sapiens

<220>

<221> TrCP(261-569)antisense

<222> (1)..(42)

<223> Beta-TrCP primer

<400> 15
gcgagcgtcg acctagggcc cttatctgga gatgtaggtg ta

42

<210> 16

<211> 36

<212> DNA

<213> homo sapiens

<220>

<221> TrCP(1-333)sense

<222> (1)..(36)

<223> Beta-TrCP primer

<400> 16
gcgagcggat ccatggaccc ggccgaggcg gtgctg

36

<210> 17

<211> 45

<212> DNA

<213> homo sapiens

<220>

<221> TrCP(1-333)antisense

<222> (1)..(45)

<223> Beta-TrCP primer

<400> 17
gcgagcgtcg acctagggcc catccacac tctgaccgtg gaatc

45

<210> 18

<211> 36

<212> DNA

<213> Homo sapiens

<220>

<221> TrCP(deltaN)(144-569)sense

<222> (1)..(36)

<223> Beta-TrCP (deltaN) primer

<400> 18

gcgagcggat ccatgttcat aactgctctg ccagct

36

<210> 19

<211> 42

<212> DNA

<213> Homo sapiens

<220>

<221> TrCP(deltaN)(144-569)antisense

<222> (1)..(42)

<223> Beta-TrCP(deltaN) primer

<400> 19

gcgagcgtcg acctagggcc cttatctgga gatgtaggtg ta

42

<210> 20

<211> 19

<212> RNA

<213> Homo sapiens

<220>

<221> RassF1dsRNA185

<222> (1)..(19)

<223> Sense strand sequence
Antisense strand SEQ ID NO.21

<220>

<221> RasSF1dsRNA185

<222> (1)..(19)

<223> Sense strand sequence (dTdT in 3')

<400> 20
gcugagauug agcagaaga

19

<210> 21

<211> 19

<212> RNA

<213> Homo sapiens

<220>

<221> RasSF1dsRNA185

<222> (1)..(19)

<223> Antisense strand (dTdT in 3')

<400> 21
cgacucuaac ucgucuucu

19

<210> 22

<211> 21

<212> RNA

<213> Homo sapiens

<220>

<221> RasSF1sequence

<222> (1)..(21)

<223> Target of Ras SF1 dsRNA 185

<400> 22
aagcugagau ugagcagaag a

21

<210> 23

<211> 19

<212> RNA

<213> Homo sapiens

<220>

<221> RasSF1dsRNA202

<222> (1)..(19)

<223> Sense strand (dTdT in 3')

<400> 23
gaucaaggag uacaaugcc

19

<210> 24

<211> 19

<212> RNA

<213> Homo sapiens

<220>

<221> RasSF1dsRNA202

<222> (1)..(19)

<223> Antisense strand (dTdT in 3')

<400> 24
cuaguuccuc auguuacgg

19

<210> 25

<211> 21

<212> RNA

<213> homo sapiens

<220>

<221> RasSF1sequence

<222> (1)..(21)

<223> target of the Ras SF1 ds RNA202

<400> 25
aagaucaagg aguacaauGC c

21

<210> 26

<211> 19

<212> RNA

<213> mammalian

<220>

<221> dsRNALucGL2

<222> (1)..(19)

<223> sense strand (dTdT in 3')

<400> 26
cguacgcgga auacuucga

19

<210> 27

<211> 19

<212> RNA

<213> Mammalian

<220>

<221> dsRNALucGL2

<222> (1)..(19)

<223> antisense strand (dTdT in 3')

<400> 27
gcaugcgccu uaugaagcu

19

<210> 28

<211> 69

<212> DNA

<213> Artificial sequence

<220>

<223> oligonucleotide used to construct pP6 plasmid

<220>

<221> oligonucleotide

<222> (1)..(69)

<223> 5'-3' single stranded

B4717A-US.ST25.txt
Complementary to Sequence ID NO.7

<400> 28
ctagccatgg ccgcaggggc cgcggccgca ctagtgggga tccttaatta aagggccact 60
ggggccccc 69

<210> 29

<211> 64

<212> DNA

<213> Artificial sequence

<220>

<223> oligonucleotide used to construct pB6 plasmid

<220>

<221> oligonucleotide

<222> (1)..(64)

<223> 5'-3' single stranded
Complementary to SEQ ID NO. 8

<400> 29
catggccgga cgggccgcgg ccgcactagt ggggatcctt aattaaagg ccactggggc 60
cccc 64

<210> 30

<211> 76

<212> DNA

<213> Artificial sequence

<220>

<223> oligonucleotide used to construct pB20 plasmid

<220>

<221> oligonucleotide

<222> (1)..(76)

<223> 5'-3' single stranded
Complementary to SEQ ID NO.9

<400> 30

B4717A-US.ST25.txt

aattcggggc cggacgggcc gcggccgcac tagtggggat ccttaattaa gggccactgg

60

ggcccctcga cctgca

76